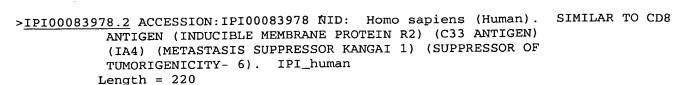
Texicoli DLASI Server

EXHIBIT "C"



Score = 451 bits (1160), Expect = e-126 Identities = 216/216 (100%), Positives = 216/216 (100%)

- Query: 33 AWLLLDRNNFLTAFDENNHFIVPISQILIGMGSSTVLFCLLGYIGIHNEIRWLLIVYAVL 92
 AWLLLDRNNFLTAFDENNHFIVPISQILIGMGSSTVLFCLLGYIGIHNEIRWLLIVYAVL 64

 Sbjct: 5 AWLLLDRNNFLTAFDENNHFIVPISQILIGMGSSTVLFCLLGYIGIHNEIRWLLIVYAVL 64

 Query: 93 ITWTFAVQVVLSAFIITKKEEVQQLWHDKIDFVISEYGSKDKPEDITKWTILNALQKTLQ 152
 ITWTFAVQVVLSAFIITKKEEVQQLWHDKIDFVISEYGSKDKPEDITKWTILNALQKTLQ 124

 Query: 153 CCGQHNYTDWIKNKNKENSGQVPCSCTKSTLRKWFCDEPLNATYLEGCENKISAWYNVNV 212
 CCGQHNYTDWIKNKNKENSGQVPCSCTKSTLRKWFCDEPLNATYLEGCENKISAWYNVNV 184
- Query: 213 LTLIGINFGLLTSEVFQVSLTVCFFKNIKNIIHAEM 248 LTLIGINFGLLTSEVFQVSLTVCFFKNIKNIIHAEM Sbjct: 185 LTLIGINFGLLTSEVFQVSLTVCFFKNIKNIIHAEM 220

Levicon Dragi Sciver

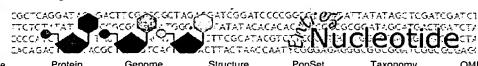
EXHIBIT "D"

>XM_084868 ACCESSION:XM_084868 NID: gi 22059998 ref XM_084868.5 Homo sapiens similar to CD82 antigen (Inducible membrane protein R2) (C33 antigen) (IA4) (Metastasis suppressor Kangai 1) (Suppressor of tumorigenicity-6) (LOC144448), mRNA Length = 992

Score = 1465 bits (739), Expect = 0.0
Identities = 746/747 (99%), Gaps = 1/747 (0%)
Strand = Plus / Plus

	Query: Sbjct:		atgttaagaaataacaaaacaataattattaagtactttcttaatctcattaatggagct	
	Query:		ttcttqqttcttggacttttattcatgggatttggtgcatggctcttattagataga	
	Sbjct:	138		196
			aatttttaacagcttttgatgaaaataatcacttcatagtacctatttctcaaattttg	
	-		aatttttaacagcttttgatgaaaataatcacttcatagtacctatttctcaaattttg attggaatgggatcttctactgttcttttttgtctattgggttatataggaattcacaac	
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-	Query:	241	gaaatcagatggctcctaattgtgtatgcagtattgataacatggacctttgctgttcag	300
•			gaaatcagatggctcctaattgtgtatgcagtattgataacatggacctttgctgttcag	
	_		gttgtactttcagcattcatcatcacaaagaaagaggggttcagcaactatggcatgac	
	Query:	361	aaaattgattttgtcatttctgagtatggatctaaagataagcctgaagatataaccaag	420
	Sbjct:	437	aaaattgattttgtcatttctgagtatggatctaaagataagcctgaagatataaccaag	496
			tggactattctgaatgccttacagaaaacattacagtgttgtggccaacataattacaca	
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PubMed Nucleotide Protein Genome **PopSet** Taxonomy **OMIM** Go Clear Search Nucleotide ▼ for Limits Preview/Index Clipboard History Details default -Display Save Text Add to Clipboard Get Subsequence

☐ 1. XM_084868. Homo sapiens simi...[gi:22059998]

Links

LOCUS LOC144448 992 bp mRNA linear PRI 01-AUG-2002 DEFINITION Homo sapiens similar to CD82 antigen (Inducible membrane protein

R2) (C33 antigen) (IA4) (Metastasis suppressor Kangai 1)

(Suppressor of tumorigenicity-6) (LOC144448), mRNA.

ACCESSION XM_084868

VERSION XM_084868.5 GI:22059998

KEYWORDS

SOURCE human.

ORGANISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (bases 1 to 992)

AUTHORS NCBI Annotation Project.

TITLE Direct Submission

JOURNAL Submitted (31-JUL-2002) National Center for Biotechnology

Information, NIH, Bethesda, MD 20894, USA

COMMENT GENOME ANNOTATION REFSEQ: This model reference sequence was

predicted from NCBI contig $\underline{\text{NT}}\underline{009538}$ by automated computational analysis using gene prediction method: BLAST, supported by EST

e<u>vidence</u>. Also see:

<u>Documentation</u> of NCBI's Annotation Process

On Aug 1, 2002 this sequence version replaced gi: 20553886.

FEATURES

Location/Qualifiers

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/chromosome="12"

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Kangai 1) (Suppressor of tumorigenicity-6) "

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EM"

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Revised: July 5, 2002.

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Sep 30 2002 14:02:06